OIPE

RAW SEQUENCE LISTING DA PATENT APPLICATION: US/09/610,313 TI

DATE: 07/17/2000 TIME: 11:06:00

Input Set : A:\seqlist.txt

Output Set: N:\CRF3\07172000\1610313.raw

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3 <110> APPLICANT: Barnett, Susan
              Zur Megede, Jan
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              POLYPEPTIDES, POLYPEPTIDES AND USES THEREOF
      9 <130> FILE REFERENCE: PP01631.101
C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/610,313
C--> 12 <141> CURRENT FILING DATE: 2000-07-05
     14 <150> PRIOR APPLICATION NUMBER: 09/475,704
     15 <151> PRIOR FILING DATE: 1999-12-30
     17 <160> NUMBER OF SEQ ID NOS: 45
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    49 ctggagaagt tegecetgaa eeceggeetg etggagacea gegagggetg caageagate 180
    50 atccgccage tgcaccccgc cetgcagace ggcagegagg agetgaagag cetgttcaac 240
    51 accytygcca coctytacty cytycacyay aagatcyayy teegegacae caaggaggee 300
    52 ctggacaaga tcgaggagga gcagaacaag tgccagcaga agatccagca ggccgaggcc 360
    53 gccgacaagg gcaaggtgag ccagaactac cccatcgtgc agaacctgca gggccagatg 420
    54 gtgcaccagg ccatcagccc ccgcaccctg aacgcctggg tgaaggtgat cgaggagaag 480
    55 gccttcagcc ccgaggtgat ccccatgttc accgccctga gcgagggcgc caccccccag 540
    56 gacctgaaca cgatgttgaa caccgtgggc ggccaccagg ccgccatgca gatgctgaag 600
    57 gacaccatca acgaggagge egeegagtgg gacegegtge acccegtgea egeeggeece 660
    58 ategececeg gecagatgeg egageceege ggeagegaca tegeeggeae caccageace 720
    59 ctgcaggagc agategeetg gatgaccage aacceeecca teeecgtggg eqacatetae 780
    60 aageggtgga teateetggg eetgaacaag ategtgegga tgtacageee egtgageate 840
    61 ctggacatca agcagggccc caaggagccc ttccgcgact acgtggaccg cttcttcaag 900
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65 ctggccgagg cgatgagcca ggccaacacc agcgtgatga tgcagaagag caacttcaag 1140
66 ggccccggc gcatcgtcaa gtgcttcaac tgcggcaagg agggccacat cgcccgcaac 1200
67 tgccgcgccc cccgcaagaa gggctgctgg aagtgcggca aggagggcca ccagatgaag 1260
68 gactgcaccg agegccaggc caactteetg ggcaagatet ggcccageca caagggccgc 1320
69 cocggeaact teetgeagag ecgeeegag eccaeegeee ecceegeega gagetteege 1380
70 ttcgaggaga ccacccccgg ccagaagcag gagagcaagg accgcgagac cctgaccagc 1440
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75 <212> TYPE: DNA
76 <213> ORGANISM: Artificial Sequence
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86 atgaageage tgeageeege eetgeagaee ggeaeegagg agetgegeag eetgtacaae 240
 87 acceptagcea coetstacts cotscaces generates techniques caassage 300
 88 ctggacaaga tcgaggagga gcagaacaag tcccagcaga agacccagca ggccaaggag 360
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 90 caccaggeca teagececeg caccetgaae geetgggtga aggtgatega ggagaaggee 480
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 92 ctgaacacga tgttgaacac cgtgggcggc caccaggccg ccatgcagat gctgaaggac 600
 93 accatcaacg aggaggccgc cgagtgggac cgcctgcacc ccgtgcaggc cggccccgtg 660
 94 geoceeggee agatgegega ecceegegge agegacateg ceggegeeae cageaceetg 720
 95 caggagcaga togoctggat gaccagcaac occoccgtgo cogtgggoga catotacaag 780
 96 eggtggatca teetgggeet gaacaagate gtgeggatgt acageceegt gageateetg 840
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 100 gaggagatga tgaccgcctg ccagggcgtg ggcggccccg gccacaaggc ccgcgtgctg 1080
 101 gccgaggcga tgagccaggc caacagcgtg aacatcatga tgcagaagag caacttcaag 1140
  102 ggcccccggc gcaacgtcaa gtgcttcaac tgcggcaagg agggccacat cgccaagaac 1200
  103 tgccgcgccc cccgcaagaa gggctgctgg aagtgcggca aggagggcca ccagatgaag 1260
  104 gactgcaceg agegccagge caactteetg ggcaagatet ggcccageca caagggcege 1320
  105 cccggcaact tcctgcagaa ccgcagcgag cccgccgccc ccaccgtgcc caccgccccc 1380
  106 cccgccgaga gcttccgctt cgaggagacc acccccgccc ccaagcagga gcccaaggac 1440
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  108 agccagtaa
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  112 <212> TYPE: DNA
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121 gccatgtacg cccccccat cgccggcaac ctgacctgcg agagcaacat caccggcctg 120
122 ctgctgaccc gcgacggcgg c
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125 <211> LENGTH: 1431
126 <212> TYPE: DNA
127 <213> ORGANISM: Artificial Sequence
130 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic
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134 agcgtggtgg gcaacctgtg ggtgaccgtg tactacggcg tgcccgtgtg gaaggaggcc 60
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137 gtgaccgaga acttcaacat gtggaagaac gacatggtgg accagatgca cgaggacatc 240
138 atcagcctgt gggaccagag cctgaagecc tgcgtgaage tgaccccct gtgcgtgacc 300
139 ctgaagtgcc gcaacgtgaa cgccaccaac aacatcaaca gcatgatcga caacagcaac 360
139 ctgaagtgee geaacgtgaa cyceaceaac aacateaaca geatgatega cacaagcaag 420
140 aagggcgaga tgaagaactg cagettcaac gtgaccaccg agetggcga cacagcaag 420
141 gaggtgeacg cectgtteta cegeetggae gtggtgeee tgeagggcaa caacagcaac 480
142 gagtaccgee tgateaactg caacaccage gecateaceac aggeetgee caaggtgage 540
143 ttegacccca tccccatcca ctactgcacc cccgccggct acgccatcct gaagtgcaac 600
 144 aaccagacet tcaacggcae eggeeeetge aacaacgtga geagegtgea gtgegeeeae 660
 145 ggcatcaagc ccgtggtgag cacccagctg ctgctgaacg gcagcctggc caagggcgag 720
 146 atcatcatcc gcagcgagaa cctggccaac aacgccaaga tcatcatcgt gcagctgaac 780
 147 aagcccgtga agatcgtgtg cgtgcgccc aacaacaaca cccgcaagag cgtgcgcatc 840
 148 ggccccggcc agacetteta egccaccggc gagateateg gcgacatecg ecaggeetae 900
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 151 accacccaca gettcaactg cegeggegag ttettetact gegacaccag ceagetgtte 1080
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 154 geoccccca tegeoggeaa ectgacetge gagageaaca teaceggeet getgetgace 1260
 155 egegacggeg geaagacegg ecceaacgae accgagatet teegeceegg eggeggegae 1320
 156 atgcgcgaca actggcgcaa cgagctgtac aagtacaagg tggtggagat caagcccctg 1380
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170 aagaccaccc tgttctgcac cagcgacgcc aaggcctacg agaccgaggt gcacaacgtg 120
  168 <400> SEQUENCE: 7
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  172 gtgaccgaga acttcaacat gtggaagaac gacatggtgg accagatgca cgaggacatc 240
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175 aagggcgaga tgaagaactg cagcttcaac gtgaccaccg agctgcgcga ccgcaagcag 420
176 gaggtgcacg ccctgttcta ccgcctggac gtggtgcccc tgcagggcaa caacagcaac 480
177 gagtaccgcc tgatcaactg caacaccagc gccatcaccc aggcctgccc caaggtgage 540
178 ttegaccea tecceateca etactgeace ecegeegget acgecatect gaagtgeaac 600
179 aaccagacct tcaacggcac cggcccctgc aacaacgtga gcagcgtgca gtgcgcccac 660
180 ggcatcaagc ccgtggtgag cacccagctg ctgctgaacg gcagcctggc caagggcgag 720
181 atcatcatcc gcagcgagaa cctggccaac aacgccaaga tcatcatcgt gcagctgaac 780
182 aagcccgtga agatcgtgtg cgtgcgccc aacaacaaca cccgcaagag cgtgcgcatc 840
183 ggccccggcc agaccttcta cgccaccggc gagatcatcg gcgacatccg ccaggcctac 900
184 tgcatcatca acaagaccga gtggaacagc accctgcagg gcgtgagcaa gaagctggag 960
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186 accacccaca getteaactg eegeggegag ttettetact gegacaccag eeagetgtte 1080
187 aacagcacct acagccccag cttcaacggc accgagaaca agctgaacgg caccatcacc 1140
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 199 atcaacact acaccgacac catctaccgc ctgctggagg agagccagaa ccagcaggag 1860
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  215 tgggccaccc acgcctgcgt gcccaccgac cccaacccc aggagatcgt gctggagaac 180
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  218 etgaagtgee gcaacgtgaa egccaccaac aacatcaaca gcatgatega caacagcaac 360
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  221 gagtaccgcc tgatcaactg caacaccagc gccatcaccc aggcctgccc caaggtgagc 540
  222 ttegacceca tececateca etactgeace ecegeegget acgecatect gaagtgeaac 600
  223 aaccagacet teaacggeae eggeeeetge aacaacgtga geagegtgea gtgegeeeae 660
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228 tgcatcatca acaagaccga gtggaacagc accctgcagg gcgtgagcaa gaagctggag 960
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233 geoccecca tegeoggeaa cetgacetge gagageaaca teaceggeet getgetgace 1260
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 269 gtgcccgtgt ggaaggaggc caagaccacc ctgttctgca ccagcgacgc caaggcctac 180
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L:11 M:270 C: Current Application Number differs, Replaced Application Number L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date